

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/943,664

DATE: 10/26/2001  
TIME: 09:32:36

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Output Set: N:\CRF3\10262001\I943664.raw

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1 <110> APPLICANT: Baker, Kevin  
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3 Eaton, Dan  
4 Ferrara, Napoleone  
5 Filvaroff, Ellen  
6 Gerritsen, Mary  
7 Goddard, Audrey  
8 Godowski, Paul  
9 Grimaldi, Christopher  
10 Gurney, Austin  
11 Hillan, Kenneth  
12 Kljavin, Ivar  
13 Napier, Mary  
14 Roy, Margaret  
15 Tumas, Daniel  
16 Wood, William  
17 <120> TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC  
18 ACIDS ENCODING THE SAME  
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Input Set : N:\Crf3\temp\09943664.raw.txt  
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100 &lt;210&gt; SEQ ID NO: 1

101 &lt;211&gt; LENGTH: 2454

102 &lt;212&gt; TYPE: DNA

103 &lt;213&gt; ORGANISM: Homo Sapien

104 &lt;400&gt; SEQUENCE: 1

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107      ctcatctttt cttcttacac agtgtctgag aacattttaca ttatagataa 150
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112      gatgatgatg atgatgatga tgatgatgat gatgaggaca actctctttt 400
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114      tgtgtccatt tggatgtcag tgctattcac gagttgtaca ttgctcagat 500
115      ttaggtttga cctcagtcac aaccaacatt ccatttgata ctggaatgct 550
116      tgatcttcaa aacaataaaa ttaaggaaat caaagaaaat gattttaaag 600
117      gactcacttc actttatggt ctgactctga acaacaacaa gctaacgaag 650
118      attcacccaa aagcctttct aaccacaaag aagttgcgaa ggctgtatct 700
119      gtcccacaat caactaagtg aaataccact taatcttccc aaatcattag 750
120      cagaactcag aattcatgaa aataaagtta agaaaataca aaaggacaca 800
121      ttcaaaggaa tgaatgcttt acacgttttg gaaatgagtg caaacctctc 850
122      tgataataat gggatagagc caggggcatt tgaaggggtg acgggtgttc 900
123      atatcagaat tgcagaagca aaactgacct cagttcctaa aggcttacca 950
124      ccaactttat tggagcttca cttagattat aataaaattt caacagtgga 1000
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126      acaacaaaat cacagatata gaaaatggga gtcttgctaa cataccacgt 1100
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135      actgacttat tttatgacaa gaaatttcaa cggaattttg ccaaactatt 1550
136      gatacataag ggggtgagag aaacaagcat ctattgcagt ttcctttttg 1600
137      cgtacaaatg atcttacata aatctcatgc ttgaccattc ctttcttcat 1650
138      aacaaaaaag taagatatcc ggtatttaac actttgttat caagcacatt 1700
139      ttaaaaagaa ctgtactgta aatggaatgc ttgacttagc aaaatttggtg 1750
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147   cgcatttttaa tgatccgcta ttataagctt ttaatagcat gaaaattggt 2150
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149   cctttggaag accttgcttg gaagagcctg gacactaaca attctacacc 2250
150   aaattgtctc ttcaaatacg tatggactgg ataactctga gaaacacatc 2300
151   tagtataact gaataagcag agcatcaaat taaacagaca gaaaccgaaa 2350
152   gctctatata aatgctcaga gttctttatg tatttcttat tggcattcaa 2400
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154   aaat 2454
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157 <211> LENGTH: 379
158 <212> TYPE: PRT
159 <213> ORGANISM: Homo Sapien
160 <400> SEQUENCE: 2
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163   Lys Pro Phe Phe Ser Pro Ser His Ile Ala Leu Lys Asn Met Met
164       20           25           30
165   Leu Lys Asp Met Glu Asp Thr Asp Asp Asp Asp Asp Asp Asp
166       35           40           45
167   Asp Asp Asp Asp Asp Glu Asp Asn Ser Leu Phe Pro Thr Arg Glu
168       50           55           60
169   Pro Arg Ser His Phe Phe Pro Phe Asp Leu Phe Pro Met Cys Pro
170       65           70           75
171   Phe Gly Cys Gln Cys Tyr Ser Arg Val Val His Cys Ser Asp Leu
172       80           85           90
173   Gly Leu Thr Ser Val Pro Thr Asn Ile Pro Phe Asp Thr Arg Met
174       95          100          105
175   Leu Asp Leu Gln Asn Asn Lys Ile Lys Glu Ile Lys Glu Asn Asp
176       110          115          120
177   Phe Lys Gly Leu Thr Ser Leu Tyr Gly Leu Ile Leu Asn Asn Asn
178       125          130          135
179   Lys Leu Thr Lys Ile His Pro Lys Ala Phe Leu Thr Thr Lys Lys
180       140          145          150
181   Leu Arg Arg Leu Tyr Leu Ser His Asn Gln Leu Ser Glu Ile Pro
182       155          160          165
183   Leu Asn Leu Pro Lys Ser Leu Ala Glu Leu Arg Ile His Glu Asn
184       170          175          180
185   Lys Val Lys Lys Ile Gln Lys Asp Thr Phe Lys Gly Met Asn Ala
186       185          190          195
187   Leu His Val Leu Glu Met Ser Ala Asn Pro Leu Asp Asn Asn Gly
188       200          205          210
189   Ile Glu Pro Gly Ala Phe Glu Gly Val Thr Val Phe His Ile Arg
190       215          220          225
191   Ile Ala Glu Ala Lys Leu Thr Ser Val Pro Lys Gly Leu Pro Pro
192       230          235          240
193   Thr Leu Leu Glu Leu His Leu Asp Tyr Asn Lys Ile Ser Thr Val
194       245          250          255
195   Glu Leu Glu Asp Phe Lys Arg Tyr Lys Glu Leu Gln Arg Leu Gly
196       260          265          270

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197   Leu Gly Asn Asn Lys Ile Thr Asp Ile Glu Asn Gly Ser Leu Ala
198                               275                280                285
199   Asn Ile Pro Arg Val Arg Glu Ile His Leu Glu Asn Asn Lys Leu
200                               290                295                300
201   Lys Lys Ile Pro Ser Gly Leu Pro Glu Leu Lys Tyr Leu Gln Ile
202                               305                310                315
203   Ile Phe Leu His Ser Asn Ser Ile Ala Arg Val Gly Val Asn Asp
204                               320                325                330
205   Phe Cys Pro Thr Val Pro Lys Met Lys Lys Ser Leu Tyr Ser Ala
206                               335                340                345
207   Ile Ser Leu Phe Asn Asn Pro Val Lys Tyr Trp Glu Met Gln Pro
208                               350                355                360
209   Ala Thr Phe Arg Cys Val Leu Ser Arg Met Ser Val Gln Leu Gly
210                               365                370                375
211   Asn Phe Gly Met
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214 <211> LENGTH: 20
215 <212> TYPE: DNA
216 <213> ORGANISM: Artificial Sequence
217 <220> FEATURE:
218 <223> OTHER INFORMATION: Synthetic Oligonucleotide Probe
219 <400> SEQUENCE: 3
220   ggaaatgagt gcaaaccctc 20
222 <210> SEQ ID NO: 4
223 <211> LENGTH: 24
224 <212> TYPE: DNA
225 <213> ORGANISM: Artificial Sequence
226 <220> FEATURE:
227 <223> OTHER INFORMATION: Synthetic Oligonucleotide Probe
228 <400> SEQUENCE: 4
229   tcccaagctg aacactcatt ctgc 24
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232 <211> LENGTH: 50
233 <212> TYPE: DNA
234 <213> ORGANISM: Artificial Sequence
235 <220> FEATURE:
236 <223> OTHER INFORMATION: Synthetic Oligonucleotide Probe
237 <400> SEQUENCE: 5
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241 <211> LENGTH: 3441
242 <212> TYPE: DNA
243 <213> ORGANISM: Homo Sapien
244 <400> SEQUENCE: 6
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247   cctccctccc tctcccccag ctgtcccgtt cgcgtcatgc cgagcctccc 150
248   ggccccgcgc gcccgctgc tgctcctcgg gctgctgctg ctcggtctcc 200
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## VERIFICATION SUMMARY

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